Applicant: Shigeaki Kato et al. Attorney's Docket No.: 14875-054001 / C1-901PCT-US

Serial No.: 09/489,198 Filed: January 20, 2000

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REMARKS

Claims 8-9 are canceled without prejudice or disclaimer. Applicants reserve the right to pursue the canceled claims in one or more continuing applications.

Claims 10 and 11 have been amended to replace the term "gene" with "nucleic acid." New dependent claims 28-45 have been added. The amended and new claims are supported throughout the application, e.g., at page 13, lines 5-6 and 14-15; and page 16, lines 1-2. Upon entry of this amendment, claims 1-7 and 10-37 will be pending and claims 10, 11 and 28-45 will be under examination.

Rejections Under 35 U.S.C. §112, First Paragraph

Enablement

Claims 8-11 are rejected because, according to the Examiner,

the specification, while being enabling for a method for screening for a nucleic acid which encodes a polypeptide that converts an inactive form of vitamin D3 into an active form, does not reasonably provide enablement for a method for screening for a gene encoding a polypeptide that converts an inactive form of vitamin D3 to an active form, or a method for screening for a gene encoding an [sic, a] polypeptide that converts a ligand precursor into a ligand. (Office action, paragraph bridging pages 2-3.)

Claims 8 and 9 are canceled and claims 10 and 11 have been amended to recite screening a test "nucleic acid" that encodes a polypeptide rather than a test "gene" that encodes a polypeptide. This amendment is implicitly supported throughout the entire application. The present claims are thus commensurate with the Examiner's acknowledged scope of enablement. Accordingly, Applicants respectfully request that the rejection be withdrawn.

Written Description

Claims 8-11 are rejected as "containing subject matter which was not described in the specification in such a way as to reasonably convey to one skilled in the relevant art that that the inventor(s), at the time the application was filed, had possession of the claimed invention." The Examiner states:

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The specification discloses methods for screening for a nucleic acid which encodes a polypeptide that converts an inactive form of vitamin D3 into an active form wherein the nucleic acid is from human or mouse (i.e., SEQ ID NO: 1, 2 see page 16). However, detailed information regarding the structural and functional requirements of the gene encoding the polypeptide, as well as structural and functional requirements of the encoded polypeptide itself are lacking. (Office action, page 5.)

This rejection has been addressed, in part, by canceling claims 8 and 9 and amending claims 10 and 11 to recite screening a "test nucleic acid" that encodes a polypeptide rather than a "test gene" that encodes a polypeptide. However, the Examiner's implication that there is written description for a test nucleic acid from only human and mouse is respectfully traversed. For at least the following reasons, a skilled artisan would recognize that Applicants were in possession of the full scope of "test nucleic acids" to be screened in the claimed methods.

The present claims are directed to methods of <u>screening</u>. By definition, agents to be tested in a method of screening (in this case, test nucleic acids that encode a polypeptide) are not limited to one particular structure or one particular source. Indeed, screening assays are routinely claimed in terms of specific "test compounds" where no particular compounds are disclosed in the specification, because the invention lies in the steps of the method, <u>not</u> in the identity of the compounds that can be run through the screening assay. Here, even more than the usual detail is given in the claims, in that the compound to be tested must be <u>a nucleic acid</u> encoding a polypeptide.

A skilled artisan would understand that a "test nucleic acid encoding a polypeptide" describes, e.g., nucleic acids found in conventional, art-recognized, routinely available DNA libraries, such as cDNA or genomic libraries, from various sources. See, e.g., the specification at page 13, lines 21-22, which provides: "Genes are screened from cells or cDNA libraries prepared from mRNA isolated from tissues or the like, which are expected to express an objective gene." As stated in the Guidelines for the Examination of Patent Applications Under the 35 U.S.C. §112, paragraph 1 "Written Description" Requirement (Federal Register, Vol. 66, No. 4, pages 1099-1111, Friday, January 5, 2001) (hereinafter "the Guidelines") "[t]he absence of definitions or details for well-established terms or procedures should not be the basis of a rejection under 35 U.S.C. § 112, paragraph 1, for lack of adequate written description." Prior to the priority

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date, libraries of test nucleic acids were well-established and readily available from numerous sources, including human, mouse, bovine, cat, chicken, fruit fly, monkey, gorilla, orangutan, gibbon, rabbit, rat, and yeast (see ATCC Catalogue of Recombinant DNA Materials, 3rd edition, 1993, page 9, copy enclosed). Indeed, the *raison d'etre* of such DNA libraries is to provide test nucleic acids from numerous sources to be screened for various purposes, and this much would be understood by a skilled artisan faced with the term "test nucleic acid encoding a polypeptide". Given the ready availability of test nucleic acids from numerous species, Applicants' disclosure of mouse and human test nucleic acids is fairly representative of the full scope of the term. In fact, it would not have been necessary to name <u>any</u> species of animal in order to establish that Applicants were in possession of the full scope of the invention. Accordingly, the written description requirement is satisfied.

In light of the foregoing, Applicants respectfully request that the rejection be withdrawn.

Enclosed is a Petition for Extension of Time along with the required fee. Please apply any other charges or credits to deposit account 06-1050.

Respectfully submitted,

Date:

Janis K. Fraser, Ph.D., J.D.

Reg. No. 34,819

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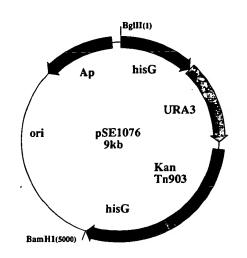
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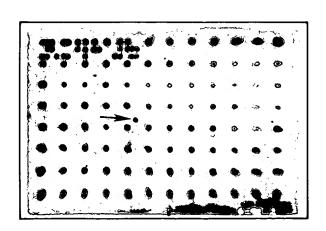
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ATCC Catalogue of Recombinant DNA Materials*

Third edition, 1993

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D.R. Maglott, Ph.D. W.C. Nierman, Ph.D.

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cDNA AND GENOMIC LIBRARIES

Name	<u> </u>]2	Library	Vector	Rest Enzyme	D١	Insert Range kb	Depositor	ATCC®	Price Code	
Alteromonas haloplanktis	g		pBR 322	Sau3A1	P	7-28 kb, 8.2 av		37436	E	1812,
Bacillus subtilis	g		Charon 4A					21.00	_	1813
bovine lymphocyte, Con A/PMA	- č			<i>Eco</i> RI	P	15 kb average	HOCH	37356	Ε	1810
stimulated, 15 hr	_		λgtll			> 0.2 kb	REEVES	37483	Ē	1777.
cat placenta	g		ILΚ						-	1816
	٥		VII	Mbol	P	17-22 kb	ROY-BURMAN	77094	Ε	3815-
chicken erythrocyte	g		λ EMBL3		_					3817
Drosophila melanogaster	g		Charon 4A	Sau3A1	P	10-20 kb	HEUMANN	37501	Ε	1964
Escherichia coli	g		λSE6			16 kb average	MANIATIS	37332	Ē	1806
_	•		ASEO	Sau3A1	P	12-17 kb	ELLEDGE.	37386	Ē	1807
Formosa monkey thymus	g	L1013	Charon 4A	<i>Eco</i> R1	_		WALKER			
gorilla lymph node	g	L1010	Charon 4A	EcoRI EcoRI	P	10-20 kb	SAKAKI	*57764	E	485
human	g	DK	Charon 4A		P	10-20 kb	SAKAKI	*57761	E	485
hu	•		Charon 4A	EcoR1	P	16-22 kb	BANK	37385	E	40,
human basal ganglia, 1-day-old	С	LMG3	λgι!I			>0010H				2731
infant, < 6 hr autolysis			· ·			> 0.8-1.0 kb	LAZZARINI	37433	Ε	116
human brain stem, I-day-old inf. < 6 hr autolysis	ant, c	LMG2	λgtH			> 0.8-1.0 kb			_	
human fetal liver			Ü			∕ 0.6-1.0 KB	LAZZARINI	37432	E	116
human leukaawa 40	g		Charon 4A	Haelll/Alul	P	15-20 kb			_	
human leukocyte, 68-year-old	g		λMG14	Mbol	P	15-20 kb	MANIATIS	37333	E	610
female, Black, with NIDDM human lymph nodes					•	13-20 KB	ROTWEIN	37458	Ε	201,
human spinol and 1 4	g	L1014	Charon 4A	Haelll/ Alul	P	15-20 kb			_	1820
human spinal cord, 1-day-old inf		LMG4	λgt!l		•	> 0.8-1.0 kb	SAKAKI	*57760	E	
human SSPE cerebellum, 7-year-	c	LMG5	λgt!!			> 0.8-1.0 kb	LAZZARINI	37434	E	116
old, < 6 hr autolysis			_			/ U.O-1.U KD	LAZZARINI	37435	E	116
human thymus, female, Caucasian, 23-year-old	c	normalized	λgt10	Eco R I	C	0.4-2 kb	_:		_	
human tonsil		thymus cDNA	•	D. O. C.		0.4-2 KD	PATANJALI,	77081	E	3779
noman folish	С	λS2T	λgtll			2-7 kb	WEISSMAN			
			•			2-7 KU	KLICKSTEIN	37546	E	284,
human tonsil										432,
manian tonsir	С	λΤ	λgtll			1.2 kb average	V.1.1044 amma	22212	_	704
mouse embryo						1.2 KU average	KLICKSTEIN	37545	E	34, 705,
mouse telencephalon, embryonic	g		Charon 28	Mbol	Р	16-20 kb	LEDER	27404	_	749
day 15	С	E171	E61	Not1	Ċ	0.3-10 kb	LEDER	37484	E	
mouse whole brain, 18-day-old					•	0.5 TO KD	RUBENSTEIN	77310	E	4479,
Onchocerca volvulus	c	LMG	λgtll			> 0.8-1.0 kb	LAZZARINI	27421		4830
Onchocerca volvulus	c		λgtil			- 010 1.0 KD	DONELSON	37431	E .	
orangutan lymphocyte	c	1 10 1 1	λZAPII			0.2-1.8 kb	DONELSON	37509 37711		2351
Pasteurella haemolytica	g	L1011	Charon 4A	EcoRI	Р	10-20 kb	SAKAKI	*57762	E E	405
	g	λEMBL4::PhBam	λEMBL4	Bam H I	С	9-25 kb	†Patent			485
Pasteurella haemolytica	_) F) (D) (D) D					deposit	40324	E	3208
and the same of th	g	λEMBL4::PhSau	λ EMBL4	Sau3A1	P	9-24 kb	†Patent	40325	E	2200
rabbit liver	•						deposit	40323	E	3208
rat brain, 2-weeks-old	g c		Charon 4A	Haelll and Alui	P	17 kb average	HARDISON	37376	E	1806
rat brain, 2-weeks-old	c		λgtll			0.5 kb average		37477		1817
rat brain, 12-weeks-old,	c		λgtll			0.3 kb average		37476		1818
cytoplasmic poly(A)+ RNA	C		λgt10			> 0.2-3 kb		37478		1814
Saccharomyces cerevisiae	g	CEN BANK	VC					3,470	L	1014
	5	CENBANK	YCp50	Sau3A1	Р	10-20 kb	ROSE	37415	E	1821.
Saccharomyces cerevisiae	o		1514515					27413		2119
Saccharomyces cerevisiae	g g		AEMBL3A	Sau3A1	P	12-15 kb	ELLEDGE	77257	E '	2117
Saccharomyces cerevisiae	-		p366	Sau3A1	P	9-12 kb			Ē	
Saccharomyces cerevisiae	g g		pRS200			8-10 kb			E	
Saccharomyces cerevisiae	g		λYES-R		S	4-8 kb				1 639
Saccharomyces cerevisiae	g		YEpi3			5-20 kb				1809
Saccharomyces cerevisiae	g	pURSCI	YRp7			5-20 kb				1809
	6	portoci	pUR18	Sau3A1	P	3-10 kb				1594
Saccharomyces cerevisiae	g	pURSC2	nliD 10	0. 1			CARR		- '	
	•	PONDEZ	pUR18	Sau3A1	P	1.5-4.0 kb		77296	E 4	1594
									- 7	
white-handed gibbon lymphocyte	g	L1012	Charon 4A	Eco R1	Р	10-20 kb	CARR			

Digest - C = complete (limit), P = partial, S = random shear.

Human chromosome-specific libraries deposited in connection with the ATCC/NIH Repository are described in the ATCC/NIH Repository Catalogue of Human and Mouse DNA Probes and Libraries.

² Insert — c = cDNA, g = genomic.

[†] This material is cited in a U.S. and/or other Patent or application and may not be used to infringe the patent claims.

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